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RAW SEQUENCE LISTING   DATE: 03/27/2003
PATENT APPLICATION: US/09/889,756A   TIME: 13:19:45

Input Set : N:\Crf4\03272003\I889756.raw
Output Set: N:\CRF4\03272003\I889756A.raw

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 1 <110> APPLICANT: Joelle Thonnard
 2 <120> TITLE OF INVENTION: BASB055
 3 <130> FILE REFERENCE: BM45353
 4 <140> CURRENT APPLICATION NUMBER: US/09/889,756A
 5 <141> CURRENT FILING DATE: 2002-04-03
 6 <150> PRIOR APPLICATION NUMBER: 9901462.3
 7 <151> PRIOR FILING DATE: 1999-01-22
 8 <150> PRIOR APPLICATION NUMBER: 9902069.5
 9 <151> PRIOR FILING DATE: 1999-01-29
10 <160> NUMBER OF SEQ ID NOS: 4
11 <170> SOFTWARE: FastSEQ for Windows Version 4.0
13 <210> SEQ ID NO: 1
14 <211> LENGTH: 1239
15 <212> TYPE: DNA
16 <213> ORGANISM: Neisseria meningitidis
17 <400> SEQUENCE: 1
18      atggctttt atgcttttaa ggcgatgcgt gcggccgcgt tggctgccgc cgttgcattg      60
19      gtactgtcgt cttgcggtaa aggcggagac gcggcgcagg gcgggcagcc tgctggtcgg     120
20      gaagcccctg cgcccgtcgt cggtgtcgta accgtccatc cgcaaaccgt cgcattgacc     180
21      gtcgagttgc cggggcgttt ggaatcgctg cgtaccgccg atgtccgcgc ccaagtcggc     240
22      ggcatcatcc aaaaacgcct gttccaagaa ggcagttatg tccgtgccgg acagccgctg     300
23      tatcagatcg acagttccac ttatgaagca aatctggaaa gcgcgcgcgc gcaactggca     360
24      acggctcagg caacgcttgc caaagcggat gcggatttgg cgcgatacaa gcctttggtt     420
25      gccgccgaag ccgtcagccg gcaggaatac gatgctgcgg taacggcgaa acgttctgcc     480
26      gaggcaggtg tcaaagcagc acaggcggca atcaaatctg ccggcattaa tctgaaccgt     540
27      tcgcgcatta ccgcgccgat ttccggcttt atcggtcagt ccaaagtttc cgaaggtacg     600
28      ctgttgaatg cgggcgatac gaccgtgctg gcaaccatcc gccaaaccaa tccgatgtat     660
29      gtgaacgtta cccagtctgc atccgaagtg atgaaattgc gccgtcagat agccgaaggc     720
30      aaactgctgg cggcggatgg tgtgattgcg gtcggcatca aatttgacga cggcacagtt     780
31      taccctgaaa aaggccgcct gctgtttgcc gatccggtcg tcaacgaatc gaccggtcag     840
32      attaccctgc gcgccgccgt accgaacgat cagaatatcc tgatgcccgg tctgtatgtg     900
33      cgcgtgctga tggaccaagt ggcggtggat aacgcatttg ttgtgccgca gcaggcggta     960
34      acgcgcggtg cgaaagatac cgtgatgatt gtgaatgccc aaggcggtat ggaaccccgc    1020
35      gaggtaacgg ttgcgcaaca gcagggtacg aattggattg ttacgtcggg tctgaaggac    1080
36      ggggacaagg tggttgtgga aggcatcagt atcgccggta aacgggtgc gaaaaaggta     1140
37      acgcccaaag aatgggcgtc gtctgaaaac caagccgccg cgcctcaatc cggcgttcag    1200
38      acggcatctg aagccaaaac tgcttctgaa gcggaataa                            1239
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 412
42 <212> TYPE: PRT
43 <213> ORGANISM: Neisseria meningitidis
44 <400> SEQUENCE: 2
45      Met Ala Phe Tyr Ala Phe Lys Ala Met Arg Ala Ala Ala Leu Ala Ala
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```
46    1              5                    10              15
47    Ala Val Ala Leu Val Leu Ser Ser Cys Gly Lys Gly Gly Asp Ala Ala
48                20                   25               30
49    Gln Gly Gly Gln Pro Ala Gly Arg Glu Ala Pro Ala Pro Val Val Gly
50                35                   40               45
51    Val Val Thr Val His Pro Gln Thr Val Ala Leu Thr Val Glu Leu Pro
52          50                   55                60
53    Gly Arg Leu Glu Ser Leu Arg Thr Ala Asp Val Arg Ala Gln Val Gly
54    65                   70                   75               80
55    Gly Ile Ile Gln Lys Arg Leu Phe Gln Glu Gly Ser Tyr Val Arg Ala
56                85                   90                   95
57    Gly Gln Pro Leu Tyr Gln Ile Asp Ser Ser Thr Tyr Glu Ala Asn Leu
58                100                  105                 110
59    Glu Ser Ala Arg Ala Gln Leu Ala Thr Ala Gln Ala Thr Leu Ala Lys
60                115                  120                 125
61    Ala Asp Ala Asp Leu Ala Arg Tyr Lys Pro Leu Val Ala Ala Glu Ala
62          130                  135                  140
63    Val Ser Arg Gln Glu Tyr Asp Ala Ala Val Thr Ala Lys Arg Ser Ala
64    145                  150                  155                 160
65    Glu Ala Gly Val Lys Ala Ala Gln Ala Ala Ile Lys Ser Ala Gly Ile
66                165                  170                 175
67    Asn Leu Asn Arg Ser Arg Ile Thr Ala Pro Ile Ser Gly Phe Ile Gly
68                180                  185                 190
69    Gln Ser Lys Val Ser Glu Gly Thr Leu Leu Asn Ala Gly Asp Thr Thr
70                195                  200                 205
71    Val Leu Ala Thr Ile Arg Gln Thr Asn Pro Met Tyr Val Asn Val Thr
72          210                  215                  220
73    Gln Ser Ala Ser Glu Val Met Lys Leu Arg Arg Gln Ile Ala Glu Gly
74    225                  230                  235                 240
75    Lys Leu Leu Ala Ala Asp Gly Val Ile Ala Val Gly Ile Lys Phe Asp
76                245                  250                 255
77    Asp Gly Thr Val Tyr Pro Glu Lys Gly Arg Leu Leu Phe Ala Asp Pro
78                260                  265                 270
79    Val Val Asn Glu Ser Thr Gly Gln Ile Thr Leu Arg Ala Ala Val Pro
80                275                  280                 285
81    Asn Asp Gln Asn Ile Leu Met Pro Gly Leu Tyr Val Arg Val Leu Met
82          290                  295                  300
83    Asp Gln Val Ala Val Asp Asn Ala Phe Val Val Pro Gln Gln Ala Val
84    305                  310                  315                 320
85    Thr Arg Gly Ala Lys Asp Thr Val Met Ile Val Asn Ala Gln Gly Gly
86                325                  330                 335
87    Met Glu Pro Arg Glu Val Thr Val Ala Gln Gln Gln Gly Thr Asn Trp
88                340                  345                 350
89    Ile Val Thr Ser Gly Leu Lys Asp Gly Asp Lys Val Val Val Glu Gly
90                355                  360                 365
91    Ile Ser Ile Ala Gly Ile Thr Gly Ala Lys Lys Val Thr Pro Lys Glu
92          370                  375                  380
93    Trp Ala Ser Ser Glu Asn Gln Ala Ala Ala Pro Gln Ser Gly Val Gln
94    385                  390                  395                 400
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 95        Thr Ala Ser Glu Ala Lys Thr Ala Ser Glu Ala Glu
 96                         405                 410
 98 <210> SEQ ID NO: 3
 99 <211> LENGTH: 41
100 <212> TYPE: DNA
101 <213> ORGANISM: Artificial Sequence
102 <220> FEATURE:
103 <223> OTHER INFORMATION: primer
104 <400> SEQUENCE: 3
105        aggcagaggc atatggcttt ttatgctttt aaggcgatgc g            41
107 <210> SEQ ID NO: 4
108 <211> LENGTH: 42
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
111 <220> FEATURE:
112 <223> OTHER INFORMATION: primer
113 <400> SEQUENCE: 4
114        aggcagaggc tcgagttccg cttcagaagc agttttggct tc          42
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VERIFICATION SUMMARY   DATE: 03/27/2003
PATENT APPLICATION: **US/09/889,756A**   TIME: 13:19:46

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